INFORMAL SEQUENCE LISTING

SEQ ID NO:1
SIZE: 10
PRT-- HIV-Tat

Tyr-Gly-Arg-Lys-Lys-Arg-Arg-Gln-Arg-Arg-Arg

SEQ ID NO: 2
SIZE: 11
PRT HIV-Tat Variant

Tyr-Ala-Arg-Lys-Ala-Arg-Arg-Gln-Ala-Arg-Arg

SEQ ID NO: 3
SIZE: 11
PRT-HIV-Tat Variant

Tyr-Ala-Arg-Ala-Ala-Ala-Arg-Gln-Ala-Arg-Ala

SEQ ID NO: 4
SIZE: 11
PRT-HIV-Tat Variant

Tyr-Ala-Arg-Ala-Ala-Arg-Ala-Ala-Arg-Arg-Arg

SEQ ID NO: 5
SIZE: 11
PRT: HIV-Tat Variant

Tyr-Ala-Arg-Ala-Ala-Arg-Ala-Ala-Arg-Arg-Ala

SEQ ID NO: 6
SIZE: 11
HIV-Tat Variant

Tyr-Ala-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg

SEQ ID NO: 7
SIZE: 11
PRT-HIV-Tat Variant

Tyr-Ala-Ala-Ala-Ala-Arg-Arg-Arg-Arg-Arg-Arg

SEQ ID NO: 8
SIZE: 11
PRT-HIV-Tat Variant

Ala-Gly-Arg-Lys-Lys-Arg-Arg-Gln-Arg-Arg-Arg

SEQ ID NO: 9
SIZE: 34
PRT-HSV VP22

Asp-Ala-Ala-Thr-Ala-Thr-Arg-Gly-Arg-Ser-Ala-Ala-Ser-Arg-Pro-Thr-Glu-Arg-
Pro-Arg-Ala-Pro-Ala-Arg-Ser-Ala-Ser-Arg-Pro-Arg-Arg-Pro-Val-Glu

SEQ ID NO: 10
SIZE: 16
PRT-Antennapedia third Helix, 43-58, Penetratin-1

Arg-Gln-Ile-Lys-Ile-Trp-Phe-Gln-Asn-Arg-Arg-Met-Lys-Trp-Lys-Lys

SEQ ID NO: 11
SIZE: 16
PRT-Antennapedia third Helix, 53-43

Lys-Lys-Trp-Lys-Met-Arg-Arg-Asn-Gln-Phe-Trp-Ile-Lys-Ile-Gln-Arg

SEQ ID NO: 12
SIZE: 16
PRT-Antennapedia third Helix, 43-58, D-amino acids

Arg-Gln-Ile-Lys-Ile-Trp-Phe-Gln-Asn-Arg-Arg-Met-Lys-Trp-Lys-Lys

SEQ ID NO: 13
SIZE: 16
PRT-Antennapedia third Helix, 43-58, Pro50,

Arg-Gln-Ile-Lys-Ile-Trp-Phe-Pro-Asn-Arg-Arg-Met-Lys-Trp-Lys-Lys

SEQ ID NO: 14
SIZE: 16
PRT-Antennapedia third Helix, 43-58, 3-Pro

Arg-Gln-Pro-Lys-Ile-Trp-Phe-Pro-Asn-Arg-Arg-Lys-Pro-Trp-Lys-Lys

SEQ ID NO: 15
SIZE:
PRT-Antennapedia third Helix, 43-58, R52M/M54R,

Arg-Gln-Ile-Lys-Ile-Trp-Phe-Gln-Asn-Met-Arg-Arg-Lys-Trp-Lys-Lys)

SEQ ID NO: 16
SIZE: 16
PRT-Antennapedia third Helix, 43-58, 7-Arg

Arg-Gln-Ile-Arg-Ile-Trp-Phe-Gln-Asn-Arg-Arg-Met-Arg -Trp-Arg -Arg

SEQ ID NO: 17
SIZE: 16
PRT-Antennapedia third Helix, 43-58, W/R

Arg-Arg-Trp-Arg-Arg-Trp-Trp-Arg-Arg-Trp-Trp-Arg-Arg-Trp-Arg-Arg

SEQ ID NO: 18
SIZE: 16
PRT-Kaposi's FGF signal sequence

Ala-Ala-Val-Ala-Leu-Leu-Pro-Ala-Val-Leu-Leu-Ala-Leu-Leu-Ala-Pro

SEQ ID NO: 19
SIZE: 20
PRT: amino terminal secretory signal of human IL-2

Met-Tyr-Arg-Met-Gln-Leu-Leu-Ser-Cys-Ile-Ala-Leu-Ser-Leu-Ala-Leu-Val-Thr-
Asn-Ser

SEQ ID NO: 20
SIZE: 20
PRT-IL-2-4 signal sequence

Met-Tyr-Arg-Met-Ala-Leu-Leu-Ser-Cys-Ile-Ala-Leu-Ser-Leu-Ala-Leu-Val-Thr-Asn-Ser

SEQ ID NO: 21
SIZE: 202
PRT-HSV VP22 sequence

Met-Thr-Ser-Arg-Arg-Ser-Val-Lys-Ser-Gly-Lys-Arg-Glu-Val-Lys-Arg-Asp-Glu-Tyr-Glu-Asp-Leu-Tyr-Tyr-Thr-Lys-Ser-Ser-Gly-Ile-Ala-Ser-Lys-Asp-Ser-Lys-Lys-Asp-Thr-Ser-Arg-Arg-Gly-Ala-Leu-Gln-Thr-Arg-Ser-Arg-Gln-Arg-Gly-Glu-Val-Arg-Phe-Val-Gln-Tyr-Asp-Glu-Ser-Asp-Tyr-Ala-Leu-Tyr-Gly-Gly-Ser-Ser-Ser-Glu-Asp-Asp-Glu-His-Pro-Glu-Val-Lys-Arg-Thr-Arg-Arg-Lys-Val-Ser-Gly-Ala-Val-Leu-Ser-Gly-Lys-Gly-Lys-Ala-Arg-Ala-Lys-Lys-Lys-Lys-Ala-Gly-Ser-Gly-Gly-Ala-Gly-Arg-Thr-Lys-Thr-Thr-Ala-Lys-Arg-Ala-Lys-Arg-Thr-Gln-Arg-Val-Ala-Thr-Lys-Ala-Lys-Ala-Ala-Lys-Ala-Ala-Glu-Thr-Thr-Arg-Gly-Arg-Lys-Ser-Ala-Gln-Lys-Glu-Ser-Ala-Ala-Leu-Lys-Asp-Ala-Lys-Ala-Ser-Thr-Ala-Lys-Thr-Arg-Ser-Lys-Thr-Lys-Ala-Gln-Gly-Leu-Ala-Arg-Lys-Leu-His-Phe-Ser-Thr-Ala-Lys-Lys-Asn-Lys-Asp-Ala-Lys-Trp-Thr-Lys-Arg-Val-Ala-Gly-Phe-Asn-Lys-Arg-Val-Phe-Cys-Ala-Ala-Val-Gly-Arg-Leu-Ala-Ala-Met-His-Ala-Arg-Met-Ala-Ala-Val-Gln-Leu-Trp-Asp-Met-Ser-Arg-Lys-Arg-Thr-Asp-Glu-Asp-Leu-Asn-Glu-Leu-Leu-Gly-Ile-Thr-Thr-Ile-Arg-Val-Thr-Val-Cys-Glu-Gly-Lys-Asn-Leu-Leu-Gln-Arg-Ala-Asn-Glu-Leu-Val-Asn-Lys-Asp-Val-Val-Gln-Asp-Val-Asp-Ala-Ala-Thr-Ala-Thr-Arg-Gly-Arg-Ser-Ala-Ala-Ser-Arg-Lys-Thr-Glu-Arg-Lys-Arg-Ala-Lys-Ala-Arg-Ser-Ala-Ser-Arg-Lys-Arg-Arg-Lys-Val-Glu-Ser

SEQ ID NO: 22
SIZE: 23
DNA-T7 RNAP promoter:

TAATACGACTCACTATAGGGAGA

SEQ ID NO: 23
SIZE 23
DNA -SP6 RNAP promoter:

ATTTAGGTGACACTATAGAAGAA

SEQ ID NO: 24
SIZE 23
DNA-T3 RNAP promoter:

AATTAACCCTCACTAAAGGGAGA

SEQ ID NO: 25
SIZE 23
DNA-K11 RNAP promoter:

AATTAGGGCACACTATAGGGAGA

SEQ ID NO:26
SIZE 24
PRT IL-4 signal sequence

Met-Gly-Leu-Thr-Ser-Gln-Leu-Leu-Pro-Pro-Leu-Phe-Phe-Leu-Leu-Ala-Cys-Ala-Gly-Asn-Phe-Val-His-Gly

SEQ ID NO:27
SIZE 302
PRT HSV VP22

Met-Thr-Ser-Arg-Arg-Ser-Val-Lys-Ser-Gly-Pro-Arg-Glu-Val-Pro-Arg-Asp-Glu-
Tyr-Glu-Asp-Leu-Tyr-Tyr-Thr-Pro-Ser-Ser-Gly-Met-Ala-Ser-Pro-Asp-Ser-Pro-
Pro-Asp-Thr-Ser-Arg-Arg-Gly-Ala-Leu-Gln-Thr-Arg-Ser-Arg-Gln-Arg-Gly-Glu-
Val-Arg-Phe-Val-Gln-Tyr-Asp-Glu-Ser-Asp-Tyr-Ala-Leu-Tyr-Gly-Gly-Ser-Ser-
Ser-Glu-Asp-Asp-Glu-His-Pro-Glu-Val-Pro-Arg-Thr-Arg-Arg-Pro-Val-Ser-Gly-
Ala-Val-Leu-Ser-Gly-Pro-Gly-Pro-Ala-Arg-Ala-Pro-Pro-Pro-Pro-Ala-Gly-Ser-
Gly-Gly-Ala-Gly-Arg-Thr-Pro-Thr-Thr-Ala-Pro-Arg-Ala-Pro-Arg-Thr-Gln-Arg-
Val-Ala-Thr-Lys-Ala-Pro-Ala-Ala-Pro-Ala-Ala-Glu-Thr-Thr-Arg-Gly-Arg-Lys-
Ser-Ala-Gln-Pro-Glu-Ser-Ala-Ala-Leu-Pro-Asp-Ala-Pro-Ala-Ser-Thr-Ala-Pro-
Thr-Arg-Ser-Lys-Thr-Pro-Ala-Gln-Gly-Leu-Ala-Arg-Lys-Leu-His-Phe-Ser-Thr-
Ala-Pro-Pro-Asn-Pro-Asp-Ala-Pro-Trp-Thr-Pro-Arg-Val-Ala-Gly-Phe-Asn-Lys-
Arg-Val-Phe-Cys-Ala-Ala-Val-Gly-Arg-Leu-Ala-Ala-Met-His-Ala-Arg-Met-Ala-
Ala-Val-Gln-Leu-Trp-Asp-Met-Ser-Arg-Pro-Arg-Thr-Asp-Glu-Asp-Leu-Asn-Glu-
Leu-Leu-Gly-Ile-Thr-Thr-Ile-Arg-Val-Thr-Val-Cys-Glu-Gly-Lys-Asn-Leu-Leu-
Gln-Arg-Ala-Asn-Glu-Leu-Val-Asn-Pro-Asp-Val-Val-Gln-Asp-Val-Asp-Ala-Ala-
Thr-Ala-Thr-Arg-Gly-Arg-Ser-Ala-Ala-Ser-Arg-Pro-Thr-Glu-Arg-Pro-Arg-Ala-
Pro-Ala-Arg-Ser-Ala-Ser-Arg-Pro-Arg-Arg-Pro-Val-Glu-Gly

SEQ ID NO:28
SIZE 6
PRT: Artificial

Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:29
SIZE 7
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:30
SIZE 8
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:31
SIZE 9
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:32
SIZE 10
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:33
SIZE 11
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:34
SIZE 12
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:35
SIZE 13
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:36
SIZE 14
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:37
SIZE 15
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:38
SIZE 16
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:39
SIZE 17
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:40
SIZE 18
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:41
SIZE 19
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:42
SIZE 20
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-
Gly-Cys

SEQ ID NO:43
SIZE 21
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-
Arg-Gly-Cys

SEQ ID NO:44
SIZE 22
PRT: Artificial

Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Arg-Gly-Cys

SEQ ID NO:45
SIZE 22
PRT: Kaposi's FGF signal sequence-full length

Met-Ser-Gly-Asp-Gly-Thr-Ala-Ala-Val-Ala-Leu-Leu-Pro-Ala-Val-Leu-Leu-Ala-Leu-Leu-Ala-Pro

SEQ ID NO:46
SIZE 10769
DNA: RO11

TGGCCATTGC ATACGTTGTA TCCATATCAT AATATGTACA TTTATATTGG CTCATGTCCA
ACATTACCGC CATGTTGACA TTGATTATTG 90
ACTAGTTATT AATAGTAATC AATTACGGGG TCATTAGTTC ATAGCCCATA TATGGAGTTC
CGCGTTACAT AACTTACGGT AAATGGCCCG 180
CCTGGCTGAC CGCCCAACGA CCCCCGCCCA TTGACGTCAA TAATGACGTA TGTTCCCATA
GTAACGCCAA TAGGGACTTT CCATTGACGT 270
CAATGGGTGG AGTATTTACG GTAAACTGCC CACTTGGCAG TACATCAAGT GTATCATATG
CCAAGTACGC CCCCTATTGA CGTCAATGAC 360
GGTAAATGGC CCGCCTGGCA TTATGCCCAG TACATGACCT TATGGGACTT TCCTACTTGG
CAGTACATCT ACGTATTAGT CATCGCTATT 450
ACCATGGTGA TGCGGTTTTG GCAGTACATC AATGGGCGTG GATAGCGGTT TGACTCACGG
GGATTTCCAA GTCTCCACCC CATTGACGTC 540
AATGGGAGTT TGTTTTGGCA CCAAAATCAA CGGGACTTTC CAAAATGTCG TAACAACTCC
GCCCCATTGA CGCAAATGGG CGGTAGGCGT 630
GTACGGTGGG AGGTCTATAT AAGCAGAGCT CGTTTAGTGA ACCGTCAGAT CGCCTGGAGA
CGCCATCCAC GCTGTTTTGA CCTCCATAGA 720
AGACACCGGG ACCGATCCAG CCTCCGCGGC CGGGAACGGT GCATTGGAAC GCGGATTCCC
CGTGCCAAGA GTGACGTAAG TACCGCCTAT 810
AGACTCTATA GGCACACCCC TTTGGCTCTT ATGCATGCTA TACTGTTTTT GGCTTGGGGC
CTATACACCC CCGCTTCCTT ATGCTATAGG 900
TGATGGTATA GCTTAGCCTA TAGGTGTGGG TTATTGACCA TTATTGACCA CTCCAACGGT
GGAGGGCAGT GTAGTCTGAG CAGTACTCGT 990
TGCTGCCGCG CGCGCCACCA GACATAATAG CTGACAGACT AACAGACTGT TCCTTTCCAT
GGGTCTTTTC TGCAGTCACC GTCGTCGACG 1080
GTATCGATAA GCTTGATCCA CCGCGGTGGC GGCCTGGCAC GACAGGTTTC CCGACTGGAA
AGCGGGCAGT GAGCGCAACG CAATTAATGT 1170
GAGTTAGCTC ACTCATTAGG CACCCCAGGC TTTACACTTT ATGCTTCCGG CTCGTATGTT
GTGTGGAATT GTGAGCGGAT AACAATTTCA 1260

CACAGGAAAC AGCTATGACC ATGATTACGC CAAGCTCCAA CGATTTAGGT GACACTATAG
AAGAGAAGGA ATTAATACGA CTCACTATAG 1350
GGAGAGAGAG AGAATTACCC TCACTAAAGG GAGGAGAAGC TTGCATGCCT GCAGGTCGAC
TCTAGAGGAT CCCCCGGGCT GCAGGAATTC 1440
CGCNNNNCCT CTCCCTCCCC CCCCCCTAAC GTTACTGGCC GAAGCCGCTT GGAATAAGGC
CGGTGTGCGT TTGTCTATAT GTTATTTTCC 1530
ACCATATTGC CGTCTTTTGG CAATGTGAGG GCCCGGAAAC CTGGCCCTGT CTTCTTGACG
AGCATTCCTA GGGGTCTTTC CCCTCTCGCC 1620
AAAGGAATGC AAGGTCTGTT GAATGTCGTG AAGGAAGCAG TTCCTCTGGA AGCTTCTTGA
AGACAAACAA CGTCTGTAGC GACCCTTTGC 1710
AGGCAGCGGA ACCCCCCACC TGGCGACAGG TGCCTCTGCG GCCAAAAGCC ACGTGTATAA
GATACACCTG CAAAGGCGGC ACAACCCCAG 1800
TGCCACGTTG TGAGTTGGAT AGTTGTGGAA AGAGTCAAAT GGCTCTCCTC AAGCGTATTC
AACAAGGGGC TGAAGGATGC CCAGAAGGTA 1890
CCCCATTGTA TGGGATCTGA TCTGGGGCCT CGGTGCACAT GCTTTACATG TGTTTAGTCG
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TCGCGAACAG TTGGCCCTTG AGCATGAGTC 2160
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GGTTGCGGAT AACGCTGCCG CCAAGCCTCT 2250
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AGCTAAGCGC GGCAAGCGCC CGACAGCCTT 2340
CCAGTTCCTG CAAGAAATCA AGCCGGAAGC CGTAGCGTAC ATCACCATTA AGACCACTCT
GGCTTGCCTA ACCAGTGCTG ACAATACAAC 2430
CGTTCAGGCT GTAGCAAGCG CAATCGGTCG GGCCATTGAG GACGAGGCTC GCTTCGGTCG
TATCCGTGAC CTTGAAGCTA AGCACTTCAA 2520
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GCAAGTTGTC GAGGCTGACA TGCTCTCTAA 2610
GGGTCTACTC GGTGGCGAGG CGTGGTCTTC GTGGCATAAG GAAGACTCTA TTCATGTAGG
AGTACGCTGC ATCGAGATGC TCATTGAGTC 2700
AACCGGAATG GTTAGCTTAC ACCGCCAAAA TGCTGGCGTA GTAGGTCAAG ACTCTGAGAC
TATCGAACTC GCACCTGAAT ACGCTGAGGC 2790
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TCCTCCTAAG CCGTGGACTG GCATTACTGG 2880
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AGCACTGATG CGCTACGAAG ACGTTTACAT 2970
GCCTGAGGTG TACAAAGCGA TTAACATTGC GCAAAACACC GCATGGAAAA TCAACAAGAA
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ACCGGAAGAC ATCGACATGA ATCCTGAGGC 3150
TCTCACCGCG TGGAAACGTG CTGCCGCTGC TGTGTACCGC AAGGACAAGG CTCGCAAGTC
TCGCCGTATC AGCCTTGAGT TCATGCTTGA 3240
GCAAGCCAAT AAGTTTGCTA ACCATAAGGC CATCTGGTTC CCTTACAACA TGGACTGGCG
CGGTCGTGTT TACGCTGTGT CAATGTTCAA 3330
CCCGCAAGGT AACGATATGA CCAAAGGACT GCTTACGCTG GCGAAAGGTA AACCAATCGG
TAAGGAAGGT TACTACTGGC TGAAAATCCA 3420
CGGTGCAAAC TGTGCGGGTG TCGATAAGGT TCCGTTCCCT GAGCGCATCA AGTTCATTGA
GGAAAACCAC GAGAACATCA TGGCTTGCGC 3510
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CCACGGCCTG AGCTATAACT GCTCCCTTCC GCTGGCGTTT GACGGGTCTT GCTCTGGCAT
CCAGCACTTC TCCGCGATGC TCCGAGATGA 3690
GGTAGGTGGT CGCGCGGTTA ACTTGCTTCC TAGTGAAACC GTTCAGGACA TCTACGGGAT
TGTTGCTAAG AAAGTCAACG AGATTCTACA 3780
AGCAGACGCA ATCAATGGGA CCGATAACGA AGTAGTTACC GTGACCGATG AGAACACTGG
TGAAATCTCT GAGAAAGTCA AGCTGGGCAC 3870
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AGTCATGACG CTGGCTTACG GGTCCAAAGA 3960
GTTCGGCTTC CGTCAACAAG TGCTGGAAGA TACCATTCAG CCAGCTATTG ATTCCGGCAA
GGGTCTGATG TTCACTCAGC CGAATCAGGC 4050
TGCTGGATAC ATGGCTAAGC TGATTTGGGA ATCTGTGAGC GTGACGGTGG TAGCTGCGGT
TGAAGCAATG AACTGGCTTA AGTCTGCTGC 4140
TAAGCTGCTG GCTGCTGAGG TCAAAGATAA GAAGACTGGA GAGATTCTTC GCAAGCGTTG
CGCTGTGCAT TGGGTAACTC CTGATGGTTT 4230
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TCAGTTCCGC TTACAGCCTA CCATTAACAC 4320
CAACAAAGAT AGCGAGATTG ATGCACACAA ACAGGAGTCT GGTATCGCTC CTAACTTTGT
ACACAGCCAA GACGGTAGCC ACCTTCGTAA 4410
GACTGTAGTG TGGGCACACG AGAAGTACGG AATCGAATCT TTTGCACTGA TTCACGACTC
CTTCGGTACC ATTCCGGCTG ACGCTGCGAA 4500
CCTGTTCAAA GCAGTGCGCG AAACTATGGT TGACACATAT GAGTCTTGTG ATGTACTGGC
TGATTTCTAC GACCAGTTCG CTGACCAGTT 4590
GCACGAGTCT CAATTGGACA AAATGCCAGC ACTTCCGGCT AAAGGTAACT TGAACCTCCG
TGACATCTTA GAGTCGGACT TCGCGTTCGC 4680
GTAACGCCAA ATCAATACGA CTCCGGATCT GAACTTGTTT ATTGCAGCTT ATAATGGTTA
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AAAAAAAAAA AACCCCCCCC CCCCCCCCCT 5040
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CAATTAATGT GAGTTAGCTC ACTCATTAGG 5130
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CTCACTATAG GGAGAGAGAG AGAATTACCC 5310
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TCCACCATAT TGCCGTCTTT TGGCAATGTG 5490
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GCCAAAGGAA TGCAAGGTCT GTTGAATGTC 5580
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TGCAGGCAGC GGAACCCCCC ACCTGGCGAC 5670
AGGTGCCTCT GCGGCCAAAA GCCACGTGTA TAAGATACAC CTGCAAAGGC GGCACAACCC
CAGTGCCACG TTGTGAGTTG GATAGTTGTG 5760
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GTACCCCATT GTATGGGATC TGATCTGGGG 5850
CCTCGGTGCA CATGCTTTAC ATGTGTTTAG TCGAGGTTAA AAAACGTCTA GGCCCCCCGA
ACCACGGGGA CGTGGTTTTC CTTTGAAAAA 5940
CACGATGATA ATATGGCCAC AACCATGGAA GACGCCAAAA ACATAAAGAA AGGCCCGGCG
CCATTCTATC CGCTGGAAGA TGGAACCGCT 6030
GGAGAGCAAC TGCATAAGGC TATGAAGAGA TACGCCCTGG TTCCTGGAAC AATTGCTTTT
ACAGATGCAC ATATCGAGGT GGACATCACT 6120
TACGCTGAGT ACTTCGAAAT GTCCGTTCGG TTGGCAGAAG CTATGAAACG ATATGGGCTG
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GCGCCCGCGA ACGACATTTA TAATGAACGT 6300
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TTGCAAAAAA TTTTGAACGT GCAAAAAAAG 6390
CTCCCAATCA TCCAAAAAAT TATTATCATG GATTCTAAAA CGGATTACCA GGGATTTCAG
TCGATGTACA CGTTCGTCAC ATCTCATCTA 6480
CCTCCCGGTT TTAATGAATA CGATTTTGTG CCAGAGTCCT TCGATAGGGA CAAGACAATT
GCACTGATCA TGAACTCCTC TGGATCTACT 6570
GGTCTGCCTA AAGGTGTCGC TCTGCCTCAT AGAACTGCCT GCGTGAGATT CTCGCATGCC
AGAGATCCTA TTTTTGGCAA TCAAATCATT 6660

CCGGATACTG CGATTTTAAG TGTTGTTCCA TTCCATCACG GTTTTGGAAT GTTTACTACA
CTCGGATATT TGATATGTGG ATTTCGAGTC 6750
GTCTTAATGT ATAGATTTGA AGAAGAGCTG TTTTTACGAT CCCTTCAGGA TTACAAAATT
CAAAGTGCGT TGCTAGTACC AACCCTATTT 6840
TCATTCTTCG CCAAAAGCAC TCTGATTGAC AAATACGATT TATCTAATTT ACACGAAATT
GCTTCTGGGG GCGCACCTCT TTCGAAAGAA 6930
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ACTGAGACTA CATCAGCTAT TCTGATTACA 7020
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GTTGTGGATC TGGATACCGG GAAAACGCTG 7110
GGCGTTAATC AGAGAGGCGA ATTATGTGTC AGAGGACCTA TGATTATGTC CGGTTATGTA
AACAATCCGG AAGCGACCAA CGCCTTGATT 7200
GACAAGGATG GATGGCTACA TTCTGGAGAC ATAGCTTACT GGGACGAAGA CGAACACTTC
TTCATAGTTG ACCGCTTGAA GTCTTTAATT 7290
AAATACAAAG GATATCAGGT GGCCCCCGCT GAATTGGAAT CGATATTGTT ACAACACCCC
AACATCTTCG ACGCGGGCGT GGCAGGTCTT 7380
CCCGACGATG ACGCCGGTGA ACTTCCCGCC GCCGTTGTTG TTTTGGAGCA CGGAAAGACG
ATGACGGAAA AAGAGATCGT GGATTACGTC 7470
GCCAGTCAAG TAACAACCGC GAAAAAGTTG CGCGGAGGAG TTGTGTTTGT GGACGAAGTA
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SEQ ID NO:47
SIZE 5742
DNA: LO53

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SEQ ID NO:48
SIZE 5313
DNA: LO59

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SEQ ID NO:49
SIZE 7940
DNA: RO23

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SEQ ID NO:51
SIZE 7002
DNA: RO71 CMV-IL2-signal sequence + first 11 amino acids of IL-2 fused to T7 RNAP

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